

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/576,051A
Source: TFWP
Date Processed by STIC: 1/19/07

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IFWP

RAW SEQUENCE LISTING

DATE: 01/19/2007

PATENT APPLICATION: US/10/576,051A

TIME: 09:59:56

Input Set : A:\PN-4-33424A-USN-Sequence-Listing-ST25-Final-Version.txt

Output Set: N:\CRF4\01192007\J576051A.raw

3 <110> APPLICANT: Novartis AG et al.
 5 <120> TITLE OF INVENTION: GENES INVOLVED IN NEURODEGENERATIVE DISORDERS
 7 <130> FILE REFERENCE: PN/4-33424A/USN
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/576,051A
 C--> 10 <141> CURRENT FILING DATE: 2006-04-18
 12 <150> PRIOR APPLICATION NUMBER: US60/513241
 13 <151> PRIOR FILING DATE: 2003-10-22
 15 <160> NUMBER OF SEQ ID NOS: 40
 17 <170> SOFTWARE: PatentIn version 3.2
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 2781
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Homo sapiens
 24 <220> FEATURE:
 25 <221> NAME/KEY: misc_feature
 26 <222> LOCATION: (546)..(546)
 27 <223> OTHER INFORMATION: n is a, c, g, or t
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 36 tttgtgcaag gcaaacctct ccagatgccca cttccaaata taggctctca ttaacaccag 240
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 40 cgcgctggcg ctgcaccaca tctggaagcc agggcgccag ggcagagacc ccggctcctg 360
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 44 tctcttgaaa atacttaaac acttttttct ctctccactg aaatctcaaa aaacagccca 480
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 54 cgatgctgta gggaccgccc tgctctcccg ccggaccgtt atccgcgccc ggcgcccgcg 780
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 60 agttactcca tgctgtcat accacaacca gagatcctca gctcaggagc atacagcccc 960
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(PJ.6)

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84 actcgaacag aatgcatttc ttcactccga agccaaatga caaataaagt ccaaaggcat 1680
86 tttctcctgt gctgaccaac caaataatat gtatagacac acacacatat gcacacacac 1740
88 acacacacac ccacagagag agagctgcaa gagcatggaa ttcattgtgt taaagataat 1800
90 cctttccatg tgaagtttaa aattactata tatttgctga tggctagatt gagagaataa 1860
92 aagacagtaa cctttctctt caaagataaa atgaaaagca cattgcatct tttcttccta 1920
94 aaaaaatgca aagatttaca ttgctgccaa atcatttcaa ctgaaaagaa cagtattgct 1980
96 ttgtaataga gtctgtaata ggatttccca taggaagaga tctgccagac gcgaactcag 2040
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114 tgaaactgag atgttgtcta tagctatgtc tataaacaac ctgaagactt gtgaaatcaa 2580
116 tgtttctttt ttaaaaaaca attttcaagt tttttttaca ataaacagtt ttgatttaaa 2640
118 atctcgtttg tatactattt tcagagactt tacttgcttc atgattagta ccaaaccact 2700
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126 <211> LENGTH: 268
127 <212> TYPE: PRT
128 <213> ORGANISM: Homo sapiens
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141 35 40 45
144 Ser Ser Gly Ala Tyr Ser Pro Ile Thr Val Trp Thr Thr Ala Ala Pro
145 50 55 60
148 Phe His Ala Gln Leu Pro Asn Gly Leu Ser Pro Leu Ser Gly Tyr Ser
149 65 70 75 80
152 Ser Ser Leu Gly Arg Val Ser Pro Pro Pro Pro Ser Asp Thr Ser Ser
153 85 90 95
156 Lys Asp His Ser Gly Ser Glu Ser Pro Ile Ser Asp Glu Glu Glu Arg
157 100 105 110
160 Leu Gln Ser Lys Leu Ser Asp Pro His Ala Ile Glu Ala Glu Lys Phe
161 115 120 125
164 Gln Cys Asn Leu Cys Asn Lys Thr Tyr Ser Thr Phe Ser Gly Leu Ala
165 130 135 140
168 Lys His Lys Gln Leu His Cys Asp Ala Gln Ser Arg Lys Ser Phe Ser
169 145 150 155 160
172 Cys Lys Tyr Cys Asp Lys Glu Tyr Val Ser Leu Gly Ala Leu Lys Met
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176 His Ile Arg Thr His Thr Leu Pro Cys Val Cys Lys Ile Cys Gly Lys
177                               180                               185                               190
180 Ala Phe Ser Arg Pro Trp Leu Leu Gln Gly His Ile Arg Thr His Thr
181                               195                               200                               205
184 Gly Glu Lys Pro Phe Ser Cys Pro His Cys Asn Arg Ala Phe Ala Asp
185                               210                               215                               220
188 Arg Ser Asn Leu Arg Ala His Leu Gln Thr His Ser Asp Val Lys Lys
189 225                               230                               235                               240
192 Tyr Gln Cys Lys Asn Cys Ser Lys Thr Phe Ser Arg Met Ser Leu Leu
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210 acacgcagcg gctcgcatcg cagagcgcag cgccggcgcg gggccgcgag aacgcagcgc      180
212 aggggagcag cccgaggcgg acaccgcgag ccgcccggca ctcccgcagt ccagccggct      240
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218 ctctgtctcc cggggcgagc gcgcagcccc gagcccgccc cgcgcctccc ggagccctcc      420
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224 tgcctggacc agcgcgctct ctggaaatag ctccggcccc ggccagggt cgtccccggc      600
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236 cttccgtacc tcaaaactag ggcggaaaag ggggaggaag tggaatgggg cgtgcatgct      960
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244 gcccttactc tgaagaatta aggagtgttt gtggggaggg ggtacagttc tgggtctagg      1200
246 aaccgaaaac caaaacattt tgctctttaa aaatctagtt agcgctcaga gagggcagga      1260
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250 gagaatgttt gtgagtgggt gttgaagagg ggggtgccgc tagaattgcg ccttggggct      1380
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262 tcggcggtga ggacaagtgc agtttcaagc cgtactccaa acccggtcgc gataagaagg      1740
264 agccgggagg cggcggtgga ggcggtggcg gtggcggggg cggcgggggg ggtgtttcgt      1800
266 cggagaagtc gggattccgg gtaccgagcg ccacctgcca gccattcacg cccaggacag      1860
268 gcagcccgag ctccagcgcc tcggcctgct cgccgggagg tatgctgtcc tcggccgggg      1920

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276 tgggctcgga ctgcggcggt tcatcgggct ccagctccgg ctccggcccc agcgcgcccc 2160
278 cctcctcctc agtgttgggc tctgggctgg tggctcccgt gtcaccctac aagccgggccc 2220
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288 gccgggaccc ttactgcctc agctaccact gcgctagcca cctggcagg gccggcgccg 2520
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306 agagcccgt tcccacgcct ggcgcccccg tgccgggtgc ccgcccacc ggaccgtact 3060
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346 ccacacgaaa aaaaaagccc cgggggggaca accagagggt tctgagagaa ccccaaaggg 4260
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350 ttgcc
4325

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353 <210> SEQ ID NO: 4

354 <211> LENGTH: 619

355 <212> TYPE: PRT

356 <213> ORGANISM: Homo sapiens

358 <400> SEQUENCE: 4

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361 1 5 10 15

364 Pro Asn Gly Glu Asn Val Cys Glu Trp Val Leu Lys Arg Gly Cys Arg

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372 Glu Ala Phe Glu Pro Gln Lys Pro Gly Phe Tyr Lys Gly Phe His Phe
373          50          55          60
376 Phe Leu Cys Val Thr Leu Pro Pro His Arg Leu Thr Pro Gln Leu Asp
377 65          70          75          80
380 Ala Lys Lys Ser Pro Leu Ala Leu Leu Ala Gln Thr Cys Ser Gln Ile
381          85          90          95
384 Gly Lys Pro Asp Pro Ser Pro Ser Ser Lys Leu Ser Ser Val Ala Ser
385          100          105          110
388 Asn Gly Gly Gly Ala Gly Gly Ala Gly Gly Gly Ala Ala Gly Asp Lys
389          115          120          125
392 Asp Thr Lys Ser Gly Pro Leu Lys Leu Ser Asp Ile Gly Val Glu Asp
393          130          135          140
396 Lys Ser Ser Phe Lys Pro Tyr Ser Lys Pro Gly Ser Asp Lys Lys Glu
397 145          150          155          160
400 Pro Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly
401          165          170          175
404 Gly Val Ser Ser Glu Lys Ser Gly Phe Arg Val Pro Ser Ala Thr Cys
405          180          185          190
408 Gln Pro Phe Thr Pro Arg Thr Gly Ser Pro Ser Ser Ser Ala Ser Ala
409          195          200          205
412 Cys Ser Pro Gly Gly Met Leu Ser Ser Ala Gly Gly Ala Pro Glu Gly
413          210          215          220
416 Lys Asp Asp Lys Lys Asp Thr Asp Val Gly Gly Gly Gly Lys Gly Thr
417 225          230          235          240
420 Gly Gly Ala Ser Ala Glu Gly Gly Pro Thr Gly Leu Ala His Gly Arg
421          245          250          255
424 Ile Ser Cys Gly Gly Gly Ile Asn Val Asp Val Asn Gln His Pro Asp
425          260          265          270
428 Gly Gly Pro Gly Gly Lys Ala Leu Gly Ser Asp Cys Gly Gly Ser Ser
429          275          280          285
432 Gly Ser Ser Ser Gly Ser Gly Pro Ser Ala Pro Thr Ser Ser Ser Val
433          290          295          300
436 Leu Gly Ser Gly Leu Val Ala Pro Val Ser Pro Tyr Lys Pro Gly Gln
437 305          310          315          320
440 Thr Val Phe Pro Leu Pro Pro Ala Gly Met Thr Tyr Pro Gly Ser Leu
441          325          330          335
444 Ala Gly Ala Tyr Ala Gly Tyr Pro Pro Gln Phe Leu Pro His Gly Val
445          340          345          350
448 Ala Leu Asp Pro Thr Lys Pro Gly Ser Leu Val Gly Ala Gln Leu Ala
449          355          360          365
452 Ala Ala Ala Ala Gly Ser Leu Gly Cys Ser Lys Pro Ala Gly Ser Ser
453          370          375          380
456 Pro Leu Ala Gly Ala Ser Pro Pro Ser Val Met Thr Ala Ser Leu Cys
457 385          390          395          400
460 Arg Asp Pro Tyr Cys Leu Ser Tyr His Cys Ala Ser His Leu Ala Gly
461          405          410          415

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RAW SEQUENCE LISTING ERROR SUMMARY
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 546
Seq#:21; N Pos. 13,20,34,53,59
Seq#:22; Xaa Pos. 3,5,10,16,18

VERIFICATION SUMMARY

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L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:48 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:540
L:2962 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:3078 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
M:341 Repeated in SeqNo=22